

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: August 15, 2002, 08:29:09 ; Search time 211.29 Seconds
(without alignments)
2681.536 Million cell updates/sec

Title: US-09-622-613a-14

Perfect score: 330
Sequence: 1 cagaactgggctacttcca.....ctggtatcggtgctgccc 330

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1736436 seqs, 858457221 residues

Total number of hits satisfying chosen parameters: 3472872

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : N_Geneseq_032802.*
1: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1980.DAT.*
2: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT.*
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15: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1994.DAT.*
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18: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1997.DAT.*
19: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1998.DAT.*
20: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1999.DAT.*
21: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2000.DAT.*
22: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2001A.DAT.*
23: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT.*
24: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	330	100.0	330	AAZ08130	Rana catesbeiana o
2	330	100.0	333	AAZ08131	Recombinant Met(-1
3	327	99.1	330	AAZ08134	Recombinant RacOR1
4	327	99.1	333	AAZ08135	Recombinant Met(-1
5	326.8	99.0	330	AAZ08132	Recombinant RacOR1
6	326.8	99.0	333	AAZ08133	Recombinant Met(-1
7	95	28.8	95	AAZ08144	PCR primer-3 for s
8	94.4	28.6	96	AAZ08148	PCR primer-2 for a
9	92.8	28.1	96	AAZ08141	PCR primer-1 for s

10	91.2	27.6	96	AAZ08147	PCR primer-1 for a
11	85	25.8	97	AAZ08139	PCR primer-1 for s
12	71.8	21.8	86	AAZ08140	PCR primer-2 for s
13	50	15.2	318	AAZ19767	Recombinant frog O
14	39.2	11.9	19205	AAZ34685	Human DNA for a no
15	39	11.8	436	ABN44186	Human breast cell
16	39	11.8	436	ABA54635	Human foetal liver
17	39	11.8	436	ABA24419	Probe #2885 for ge
18	39	11.8	436	AAK02925	Human brain expres
19	39	11.8	436	AAK28369	Human bone marrow
20	39	11.8	436	AAI12933	Probe #2866 for ge
21	39	11.8	436	AAI34296	Probe #2882 used t
22	39	11.8	436	AAI02854	R. pipiens recombi
23	38.2	11.6	315	AAAT94959	R. pipiens recombi
24	38.2	11.6	318	AAAT94958	R. pipiens recombi
25	38.2	11.6	321	AAAT94954	R. pipiens recombi
26	38.2	11.6	333	AAAT94957	R. pipiens recombi
27	38.2	11.6	336	AAAT94955	R. pipiens recombi
28	38.2	11.6	753	AAAT94972	R. pipiens recombi
29	38.2	11.6	768	AAAT94973	R. pipiens recombi
30	38.2	11.6	1065	AAAT94971	R. pipiens recombi
31	38.2	11.6	1065	AAAT94963	R. pipiens recombi
32	38.2	11.6	1065	AAAT94967	R. pipiens recombi
33	38.2	11.6	1074	AAAT94968	R. pipiens recombi
34	38.2	11.6	1098	AAAT94970	R. pipiens recombi
35	38.2	11.6	1137	AAAT94964	R. pipiens recombi
36	38	11.5	5997	AAO12188	Odonotoglossum ring
37	38	11.5	6597	AAO38106	ORSV CDNA. Odonot
38	37.6	11.4	312	AAZ08124	Rana pipiens liver
39	37.6	11.4	312	AAZ08125	Rana pipiens liver
40	37.6	11.4	312	AAZ08126	Recombinant RapLr1
41	37.6	11.4	315	AAZ08127	Recombinant Met(-1
42	37.6	11.4	315	AAZ08129	Recombinant Met(-1
43	37.6	11.4	315	AAZ08129	Recombinant Met(-1
44	37.6	11.4	2855	AAZ08136	Rana pipiens ribom
45	37.4	11.3	2574	AAV26293	Recombinant botul1

ALIGNMENTS

RESULT 1	
AAZ08130	standard; CDNA: 330 BP.
ID	AAZ08130;
XX	AAZ08130;
AC	25-JAN-2000 (first entry)
XX	
DT	Rana catesbeiana oocyte ribonuclease (RacOR1) encoding CDNA.
XX	
DE	
XX	Rana catesbeiana oocyte ribonuclease; RacOR1; covalently bound; CD22;
XX	IL2 antibody; ligand binding moiety; cancerous B cell; Kaposi's Sarcoma;
KW	human chorionic gonadotropin; hcg; recombinant ribonuclease; bullfrog;
KW	signal peptide; cytotoxic fusion protein; cancer; autoimmune disease;
KW	RNase; ss.
XX	
OS	Rana catesbeiana.
OS	Synthetic.
XX	
FH	Key
FT	mat_peptide
FT	Location/Qualifiers
FT	1..330
FT	/*tag= a
FT	/product= "RacOR1"
FT	/note= "Rana catesbeiana oocyte ribonuclease"
XX	
PN	W09950398-A2.
XX	
XX	07-OCT-1999.
PD	
XX	
XX	26-MAR-1999; 99WO-US06641.
PF	
XX	
XX	27-MAR-1998; 98US-0079751.

KM RacOR1, CD22; covalently bound; IL2 antibody; ligand binding moiety;
 KM cancerous B cell; Kapost's sarcoma; human chorionic gonadotropin; hCG;
 KM signal peptide; recombinant ribonuclease; cytotoxic fusion protein;
 KW cancer; bullfrog; RNase; autoimmune disease; ss.
 XX Rana catesbeiana.
 OS Synthetic.
 FH Key
 FH mat_peptide 1..333
 FT Location/Qualifiers
 FT /tag= a
 FT /product= "Recombinant Met(-1) RacOR1 Met22Leu Met57Leu"
 FT /note= "Rana catesbeiana oocyte ribonuclease"
 FT 1..3
 FT misc_feature
 FT /tag= b
 FT /note= "Additional ATG codon not found in native RacOR1"
 FT replace(67..69, ATG)
 FT /tag= c
 FT old_sequence
 FT /replace(172..174, ATG)
 FT /tag= d
 PN WO9950398-A2.
 XX
 PD 07-OCT-1999.
 XX
 XX 26-MAR-1999; 99WO-US06641.
 XX
 XX 27-MAR-1998; 98US-0079751.
 XX
 PA (USSH) US DEPT HEALTH & HUMAN SERVICES.
 XX
 PI Newton DL, Rybak SM;
 XX
 XX WPI: 1999-610847/52.
 DR P-PSDB; AAY28876.
 XX
 PT New recombinant ribonucleases, used for killing target cells, e.g. for
 PT treating cancers, viral infections or autoimmune diseases
 PS
 PS Disclosure: Page 65; 71pp; English.
 CC The present sequence is a cDNA encoding recombinant Rana catesbeiana
 CC oocyte ribonuclease (RacOR1) with Met at position 1, Met23Leu and
 CC Met58Leu. Carboxy terminal end of recombinant RacOR1 has a covalently
 CC bound ligand binding moiety, which can be a IL2 antibody directed against
 CC CD22 on cancerous B cells or human chorionic gonadotropin (hCG)
 CC effective against Kapost's sarcoma cells. Recombinant ribonucleases can
 CC be expressed in bacteria without an N-terminal methionine due to the
 CC presence of a signal peptide that is cleaved by bacteria. The soluble
 CC expression of ribonuclease allows the proteins to be fused in-frame with
 CC ligand binding moieties to form cytotoxic fusion proteins. They can be
 CC used for treatment of cancer and autoimmune diseases.
 XX
 XX Sequence 333 BP; 81 A; 97 C; 57 G; 98 T; 0 other;
 SQ
 Query Match 99.0%; Score 326.8; DB 20; Length 333;
 Best Local Similarity 99.4%; Pred. NO. 4.3e-91;
 Matches 328; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Db 184 tctactactcgtttccagctgaacactgacactgactctactcactcactcgcgtctgc 243
 Oy 241 ccgtactctctcgtactgaactaactacatctggttaaatgcaaacagatcccg 300
 Db 244 ccgtactctctcgtactgaactaactacatctggttaaatgcaaacagatcccg 303
 Oy 301 gtccattcgtcgtatcggtcgttcgccg 330
 Db 304 gtccattcgtcgtatcggtcgttcgccg 333
 RESULT 7
 AAZ08144
 ID AAZ08144 standard; DNA; 95 BP.
 XX
 XX AAZ08144;
 AC
 XX 25-JAN-2000 (first entry)
 XX
 XX PCR primer-3 for synthesising 3' half of RacOR1 gene.
 XX
 XX PCR primer: ribonuclease; RNase: RacOR1; Rana catesbeiana: mutation;
 KM recombinant RNase; ss.
 OS Synthetic.
 OS Rana catesbeiana.
 XX
 XX WO9950398-A2.
 XX
 XX 07-OCT-1999.
 XX
 XX 26-MAR-1999; 99WO-US06641.
 XX
 XX 27-MAR-1998; 98US-0079751.
 XX
 PA (USSH) US DEPT HEALTH & HUMAN SERVICES.
 XX
 PI Newton DL, Rybak SM;
 XX
 XX WPI: 1999-610847/52.
 DR
 XX
 XX New recombinant ribonucleases, used for killing target cells, e.g. for
 PT treating cancers, viral infections or autoimmune diseases
 PS
 PS Example 5; Page 41; 71pp; English.
 CC The present sequence is a PCR primer comprising the 3' half of
 CC ribonuclease (RacOR1) gene from Rana catesbeiana. It is used along with
 CC other primers to synthesise the 3' half of RNase with mutations resulting
 CC in recombinant RNase with Met22Leu and Met57Leu.
 XX
 XX Sequence 95 BP; 21 A; 29 C; 19 G; 26 T; 0 other;
 SQ
 Query Match 28.8%; Score 95; DB 20; Length 95;
 Best Local Similarity 100.0%; Pred. NO. 9.6e-20;
 Matches 95; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
XX 25-JAN-2000 (first entry)
XX
XX PCR primer-2 for assembling mutated RacOR1 gene.
DE
XX
XX PCR primer: assemble; Rana catesbelana ribonuclease gene; RacOR1; RNase;
XX mutated; ss.
XX
XX Synthetic.
OS Rana catesbelana.
XX
XX WO950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99MO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM;
XX
XX WPI; 1999-610847/52.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX Example 5; Page 41; 71pp; English.
XX
XX The present sequence is a PCR primer, used along with another primer
XX to assemble mutated Rana catesbelana ribonuclease (RacOR1) gene
XX resulting in RNase with Met22Leu and Met57Leu.
XX
XX Sequence 96 BP; 17 A; 32 C; 17 G; 30 T; 0 other;
SQ

Query Match      28.6%; Score 94.4; DB 20; Length 96;
Best Local Similarity 99.0%; Pred. No. 1.5e-19;
Matches 95; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 148 atctgcactggtgttatcaacatgaacgtctgtctactactcgtttccagctgaacact 207
   |||||||
DB 1 atctgcactggtgttatcaacatgaacgtctgtctactactcgtttccagctgaacact 60

OY 208 tgcactcgtactctatcactccgcgctgcgtgcgcg 243
   |||||||
DB 61 tgcactcgtactctatcactccgcgctgcgtgcgcg 96

RESULT 9
AAZ08141
ID AAZ08141 standard; DNA; 96 BP.
XX
XX AAZ08141;
AC
XX
XX 25-JAN-2000 (first entry)
DT
XX
XX PCR primer-1 for synthesizing 3' half of RacOR1 gene.
DE
XX
XX PCR primer: ribonuclease; RNase; RacOR1; Rana catesbelana; mutation;
XX recombinant RNase; ss.
XX
XX Synthetic.
OS Rana catesbelana.
XX
XX WO950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99MO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
```

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XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM;
XX
XX WPI; 1999-610847/52.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX Example 5; Page 41; 71pp; English.
XX
XX The present sequence is a PCR primer comprising the 3' half of
XX ribonuclease (RacOR1) gene from Rana catesbelana. It is used along with
XX other primers to synthesise the 3' half of RNase with mutations resulting
XX in recombinant RNase with Met22Leu and Met57Leu.
XX
XX Sequence 96 BP; 18 A; 31 C; 17 G; 30 T; 0 other;
SQ

Query Match      28.1%; Score 92.8; DB 20; Length 96;
Best Local Similarity 97.9%; Pred. No. 4.6e-19;
Matches 94; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 148 atctgcactggtgttatcaacatgaacgtctgtctactactcgtttccagctgaacact 207
   |||||||
DB 1 atctgcactggtgttatcaacatgaacgtctgtctactactcgtttccagctgaacact 60

OY 208 tgcactcgtactctatcactccgcgctgcgtgcgcg 243
   |||||||
DB 61 tgcactcgtactctatcactccgcgctgcgtgcgcg 96

RESULT 10
AAZ08147
ID AAZ08147 standard; DNA; 96 BP.
XX
XX AAZ08147;
AC
XX
XX 25-JAN-2000 (first entry)
DT
XX
XX PCR primer-1 for assembling mutated RacOR1 gene.
DE
XX
XX PCR primer: assemble; Rana catesbelana ribonuclease gene; RacOR1; RNase;
XX mutated; ss.
XX
XX Synthetic.
OS Rana catesbelana.
XX
XX WO950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99MO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM;
XX
XX WPI; 1999-610847/52.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX Example 5; Page 41; 71pp; English.
XX
XX The present sequence is a PCR primer, used along with another primer
XX to assemble mutated Rana catesbelana ribonuclease (RacOR1) gene
XX resulting in RNase with Met22Leu and Met57Leu.
XX
XX Sequence 96 BP; 30 A; 28 C; 16 G; 22 T; 0 other;
SQ
```

Query Match 27.6%; Score 91.2; DB 20; Length 96;
Best Local Similarity 96.9%; Pred. No. 1.4e-18;
Matches 93; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 cagaactgggtacttccagcagaacatcatcgaactcgatctgcaacact 60
|||||
DB 1 cagaactgggtacttccagcagaacatcatcgaactcgatctgcaacact 60
|||||

QY 61 atcatggacaacaacatcatcgttggtgtag 96
|||||
DB 61 atccctgcagaacaacatcatcgttggtgtag 96
|||||

RESULT 11
AAZ08139
ID AAZ08139 standard; DNA; 97 BP.
AC AAZ08139;
XX
XX
XX 25-JAN-2000 (first entry)
XX
XX PCR primer-1 for synthesizing 5' half of RACOR1 gene.
XX
XX PCR primer; ribonuclease; RNase; RACOR1; Rana catesbeiana; mutation;
XX
XX recombinant RNase; ss.
XX
XX Synthetic.
XX
XX Rana catesbeiana.
XX
XX
XX Key Location/Qualifiers
XX
XX misc_feature 22
XX
XX /*tag= a
XX /note= "Unknown additional base"
XX
XX W09950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99WO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
XX
XX (USSH) US DEPT HEALTH & HUMAN SERVICES.
XX
XX
XX Newton DL, Rybak SM;
XX
XX WPI: 1999-610847/52.
XX
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX
XX Example 5; Page 40; 71pp; English.
XX
XX
XX The present sequence is a PCR primer comprising the 5' half of
XX ribonuclease (RACOR1) gene from Rana catesbeiana. It is used along with
XX other primers to synthesise the 5' half of RNase with mutations resulting
XX in recombinant RNase with Met22Leu and Met57Leu.
XX
XX
XX Sequence 97 BP; 31 A; 27 C; 16 G; 22 T; 1 other;

Query Match 25.8%; Score 85; DB 20; Length 97;
Best Local Similarity 99.0%; Pred. No. 1.2e-16;
Matches 96; Conservative 0; Mismatches 0; Indels 1; Gaps 1;

QY 1 cagaactgggtacttccagcagaacatcatcgaactcgatctgcaacac 59
|||||
DB 1 cagaactgggtacttccagcagaacatcatcgaactcgatctgcaacac 60
|||||

QY 60 tatcatggacaacaacatcatcgttggtgtag 96
|||||

DB 61 tatcatggacaacaacatcatcgttggtgtag 97
|||||

RESULT 12
AAZ08140
ID AAZ08140 standard; DNA; 86 BP.
XX
XX
XX AAZ08140;
XX
XX
XX 25-JAN-2000 (first entry)
XX
XX
XX PCR primer-2 for synthesizing 5' half of RACOR1 gene.
XX
XX
XX PCR primer; ribonuclease; RACOR1; Rana catesbeiana; mutation; RNase;
XX recombinant RNase; ss.
XX
XX Synthetic.
XX
XX Rana catesbeiana.
XX
XX W09950398-A2.
XX
XX
XX 07-OCT-1999.
XX
XX
XX 26-MAR-1999; 99WO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
XX
XX (USSH) US DEPT HEALTH & HUMAN SERVICES.
XX
XX
XX Newton DL, Rybak SM;
XX
XX WPI: 1999-610847/52.
XX
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX
XX Example 5; Page 40; 71pp; English.
XX
XX
XX The present sequence is a PCR primer comprising the 5' half of
XX ribonuclease (RACOR1) gene from Rana catesbeiana. It is used along with
XX other primers to synthesise the 5' half of RNase with mutations resulting
XX in recombinant RNase with Met22Leu and Met57Leu.
XX
XX
XX Sequence 86 BP; 19 A; 20 C; 16 G; 31 T; 0 other;

Query Match 21.8%; Score 71.8; DB 20; Length 86;
Best Local Similarity 96.6%; Pred. No. 1.3e-12;
Matches 84; Conservative 0; Mismatches 2; Indels 1; Gaps 1;

QY 79 tacatcgttggtgcagcagaacgtgtaaacattcatcatctctgtctactact 138
|||||
DB 1 tacatcgttggtgcagcagaacgtgtaaacattcatcatctctgtctactact 59
|||||

QY 139 gttaaagctatctgcactggtgtatc 165
|||||
DB 60 gttaaagctatctgcactggtgtatc 86
|||||

RESULT 13
AAZ19767
ID AAZ19767 standard; CDNA; 318 BP.
XX
XX
XX AAZ19767;
XX
XX
XX 01-DEC-1999 (first entry)
XX
XX
XX Recombinant frog Onconase CDNA.
XX
XX
XX Ribonuclease; protein synthesis; inhibition; cancer; cytotoxic; ds.
XX
XX
XX Rana pipiens.
XX

FH	Key		Location/Qualifiers
FT	CDS	1..318	/+tag=
FT		a	/product= "Recombinant frog Oncogene"
PN			
MW		909646389-AI.	
PD			
XX		16-SEP-1999.	
PX			
PF		11-MAR-1999;	99WO-US04252.
PR		11-MAR-1998;	98US-0077557.
PA		(IMMUNO-) IMMUNOMEDICS INC.	
PI		Goldenberg DM, Hansen H, Leung S;	
DR		MP1: 1999-551416/46.	
DR		P-PSDB; MAY39400.	
PT		A new recombinant Oncogene used to treat, e.g. colon cancer -	
XX			
PS		Example 1; Fig 1; 42pp; English.	
CC		This sequence represents recombinant frog Oncogene cDNA. Oncogene has	
CC		ribonuclease and anti-tumour activity. The cDNA was produced via PCR	
CC		(using primers AAZ19766-219769) of two synthetic DNAs whose sequences	
CC		encoded most of the N-terminal or the C-terminal amino acids of mature	
CC		Oncogene. The two PCR products generated encoded either the N-terminal	
CC		54 amino acids (minus the initial methionine) or the C-terminal 51 amino	
CC		acids, and were ligated in frame at an XbaI site. The cDNA was then	
CC		subcloned into a vector e.g., pBluescript, where the AUG initiation	
CC		codon was ligated to the cDNA. After expression in E. coli, the	
CC		recombinant protein was purified. The initial N-formyl methionine was	
CC		cleaved off and the now N-terminal glutamate residue cyclised to form an	
CC		N-terminal pyroglutamate. The pyroglutamate residue forms part of the	
CC		phosphate binding pocket of oncogene and is essential for both	
CC		ribonuclease and anti-tumour activity. Oncogene is a 12 kd ribonuclease	
CC		which causes cell death as a result of potent inhibition of protein	
CC		synthesis by a mechanism involving inactivation of cellular RNA. It is	
CC		not inhibited by mammalian placental ribonuclease inhibitor, which may	
CC		explain its enhanced cytotoxicity relative to mammalian enzymes. It has	
CC		anti-tumour activity against a variety of solid tumours e.g. colon or	
CC		pancreatic cancers, and can be used alone or in combination with other	
CC		anti-cancer agents such as tamoxifen. When used as an anti-tumour agent,	
CC		Oncogene can be conjugated to a marker which targets it to a specific	
CC		cell type.	
XX			
SQ		Sequence 318 BP; 99 A; 65 C; 71 G; 83 T; 0 other:	
	Query Match	15.2%; Score 50; DB 20; Length 318;	
	Best Local Similarity	54.2%; Pred. No. 1.2e-05;	
	Matches 179; Conservative 0; Mismatches 130; Indels 21; Gaps		
OY	1	cagaactgagctacttccaggacaagaatacatcacacctcgcattcgcaaacat 60	
DB	4	cagagtgttgccaagtttcagagaagaatacatcacggataaac-----gagatgta 54	
OY	61	atcataggaacaacatctacatcggttggtgtgaagtcgaagaaagtgtaaaccatttcac 120	
DB	55	gacttgcagcaaatatratytlctcagaaatcgtttcaactgtaaggtaagatatcatttata 114	
OY	121	atctcttgtctaactcgtttaagcttatctgcacgttggtttac---aacatgaagctt 177	
DB	115	tacagtgcgccagagccgctaagaagctatccgttaaaagccattatcgcggtuaagaacgtg 174	
OY	178	ctgtactactcgttttcacagctgaacaactgacactgtaactcttacactccgcgtccg 237	
DB	175	ctgactactccaggtatctatctatcttgcgatgtgaatgtgacttca-----cgccccc 225	
OY	238	tgcctgactcttctctgtactggaactactacatctgcgttggttaagtcgaanaaccgtlac 297	

Db	226	tgcaataataagctgtaagaagaaacactaacaatttgcgttaacttgcgagaccaggct	285
Qy	298	ccggtcatttcgcgtgcgtatcgatcgctgtc	327
Db	286	cctgtacatttcgttggtagtgcgggagctgc	315
RESULT 14			
ID	AAS34685		
XX	AAS34685 standard; DNA; 19205 BP.		
XX	AAS34685;		
XX			
DT	17-DEC-2001 (first entry)		
XX			
DE	Human DNA for a novel foetal antigen, SEQ ID No 2109.		
XX			
KW	Human; foetal tissue antigen; ds; antiinflammatory; neuroprotective;		
KW	immunomodulator; cardiovascular; cytostatic; nephrotoxic;		
KW	cardiovascular; autoimmune disease; rheumatoid arthritis;		
KW	hyperproliferative disorder; breast neoplasm; cancer;		
KW	cardiovascular disorder; cardiac arrest; cerebrovascular disorder;		
KW	cerebral ischaemia; angiogenesis; nervous system disorder;		
KW	Alzheimer's disease; infection; ocular disorder; corneal infection;		
KW	wound healing; epithelial cell proliferation; food additive.		
XX			
OS	Homo sapiens.		
XX			
PN	WO200155312-A2.		
PD			
XX			
XX	02-AUG-2001.		
PF			
PR	17-JAN-2001; 2001WO-US01321.		
XX			
PR	31-JAN-2000; 2000US-0179065.		
PR	04-FEB-2000; 2000US-01806528.		
PR	24-FEB-2000; 2000US-0184664.		
PR	02-MAR-2000; 2000US-0186350.		
PR	16-MAR-2000; 2000US-0189874.		
PR	17-MAR-2000; 2000US-0190076.		
PR	18-APR-2000; 2000US-0198123.		
PR	19-MAY-2000; 2000US-0205515.		
PR	07-JUN-2000; 2000US-0209467.		
PR	28-JUN-2000; 2000US-0214886.		
PR	30-JUN-2000; 2000US-0215135.		
PR	07-JUL-2000; 2000US-0216647.		
PR	07-JUL-2000; 2000US-0216880.		
PR	11-JUL-2000; 2000US-0217487.		
PR	11-JUL-2000; 2000US-0217496.		
PR	14-JUL-2000; 2000US-0218290.		
PR	26-JUL-2000; 2000US-0220963.		
PR	26-JUL-2000; 2000US-0220964.		
PR	14-AUG-2000; 2000US-0224518.		
PR	14-AUG-2000; 2000US-0224519.		
PR	14-AUG-2000; 2000US-0225213.		
PR	14-AUG-2000; 2000US-0225214.		
PR	14-AUG-2000; 2000US-0225266.		
PR	14-AUG-2000; 2000US-0225267.		
PR	14-AUG-2000; 2000US-0225268.		
PR	14-AUG-2000; 2000US-0225270.		
PR	14-AUG-2000; 2000US-0225447.		
PR	14-AUG-2000; 2000US-0225757.		
PR	14-AUG-2000; 2000US-0225758.		
PR	14-AUG-2000; 2000US-0225759.		
PR	18-AUG-2000; 2000US-0226279.		
PR	22-AUG-2000; 2000US-0226681.		
PR	22-AUG-2000; 2000US-0226868.		
PR	22-AUG-2000; 2000US-0227182.		
PR	23-AUG-2000; 2000US-0227009.		
PR	30-AUG-2000; 2000US-0228924.		
PR	01-SEP-2000; 2000US-0229287.		
PR	01-SEP-2000; 2000US-0229343.		
PR	01-SEP-2000; 2000US-0229344.		


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Db      1017 ctaattatcatacattcattcgatccaacagtttctcccttaaccccttagacctcaca 1076
Qy      220 tctatcacctcgcgctgcgcgtaccctctcttcctgactgaaactaactacatctgagt 279
Db      1077 cctacactaccctctccccaataactactattatatttgaataatttactgctctttc 1136
Qy      280 aaatgcga 287
Db      1137 aactgcga 1144

RESULT 15
ID      ABA44186 standard; DNA; 436 BP.
XX
AC      ABA44186;
XX
DT      01-FEB-2002 (first entry)
XX
DE      Human breast cell single exon nucleic acid probe #2881.
XX
KM      Human: microarray: single exon probe; gene expression; breast;
KW      disease; cancer; ss.
XX
OS      Homo sapiens.
XX
PN      MO200157271-A2.
XX
PD      09-AUG-2001.
XX
PE      30-JAN-2001; 2001MO-US000662.
XX
PR      04-FEB-2000; 2000US-0180312.
PR      26-MAY-2000; 2000US-0207456.
PR      30-JUN-2000; 2000US-0608408.
PR      03-AUG-2000; 2000US-0632366.
PR      21-SEP-2000; 2000US-0234687.
PR      27-SEP-2000; 2000US-0236359.
PR      04-OCT-2000; 2000GB-0024263.
XX
XX
PA      (MOLE-) MOLECULAR DYNAMICS INC.
XX
PI      Penn SG, Hanzel DK, Chen W, Rank DR;
XX
DR      MPI; 2001-496933/54.
XX
PT      New spatially-addressable set of single exon nucleic acid probes,
PT      useful for measuring gene expression in sample derived from human
PT      breast, comprises number of single exon nucleic acid probes -
XX
PS      Claim 1; SEQ ID NO 2881; 327bp + sequence listing; English.
XX
XX
CC      The invention relates to a spatially-addressable set of single exon
CC      nucleic acid probes for measuring gene expression in a sample derived
CC      from human breast and Br 474 cells. The method involves contacting
CC      the probes with a collection of detectably labelled nucleic acids
CC      derived from mRNA of human breast, and then measuring the label
CC      bound to each probe of the microarray. The probes are useful for
CC      verifying the expression of regions of genomic DNA predicted to
CC      encode proteins. They are useful for gene discovery and for
CC      determining predisposition and/or prognosing breast disease. Gene
CC      expression analysis is useful for assessing the toxicity of chemical
CC      agents on cells. The microarray of this invention presents a far greater
CC      diversity of probes for measuring gene expression, with far less bias
CC      than expressed sequence tag microarrays. The method is suitable for
CC      rapid production of functional information from genomic sequence. The
CC      present sequence is a single exon nucleic acid probe of the invention.
CC      Note: The sequence data for this patent did not form part of the
CC      printed specification, but was obtained in electronic format directly
CC      from WIPO at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ      Sequence 436 BP; 137 A; 144 C; 15 G; 140 T; 0 other;
```

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Query Match      11.8%; Score 39; DB 22; Length 436;
Best Local Similarity 47.4%; Pred. No. 0.035;
Matches 117; Conservative 0; Mismatches 130; Indels 0; Gaps 0;

Qy      34 atcaacactccgatactatcgaacactatcatggaacaacaacatactacgttgyt 93
Db      112 atcactactaccactactactactactactactactactactactactactact 171
Qy      94 cagtgcaaaagtgtaacacttcatcactcctctgctactactactggttaagctatctgc 153
Db      172 atcattactactactaccaccctccaccattactactactactactactactact 231
Qy      154 actggtgtatcaacatgaacgttctgtctactactcgtttccagctgaacacttgcact 213
Db      232 gcttctataattactactcctactcctaccattactactactagtagtaccacttact 291
Qy      214 cgtacttctatcactccggtcgcgtgactctctcgtactgaactaactacatc 273
Db      292 actactactatcattactactatcaccaccacttactactactactactactaccacc 351
Qy      274 tgcgtta 280
Db      352 accacta 358
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